

#7



PCT09

RAW SEQUENCE LISTING                      DATE: 05/21/2002  
 PATENT APPLICATION: US/10/018,729      TIME: 15:42:53

Input Set : A:\ST99021 Sequence.ST25.txt  
 Output Set: N:\CRF3\05212002\J018729.raw

ENTERED

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3 <110> APPLICANT: DARTEIL, Raphael
4      CROUZET, Joel
5      STAELS, Bart
6      MAHFOUDI, Abderrahim
8 <120> TITLE OF INVENTION: SYSTEM OR REGULATION OF EXPRESSION USING PPAR NUCLEAR
RECEPTORS
10 <130> FILE REFERENCE: ST99021 US PCT
12 <140> CURRENT APPLICATION NUMBER: 10/018,729
C--> 13 <141> CURRENT FILING DATE: 2002-04-22
15 <150> PRIOR APPLICATION NUMBER: FR 99/07957
16 <151> PRIOR FILING DATE: 1999-06-22
18 <150> PRIOR APPLICATION NUMBER: US 60/149,721
19 <151> PRIOR FILING DATE: 1999-08-20
21 <150> PRIOR APPLICATION NUMBER: PCT/FR00/01744
22 <151> PRIOR FILING DATE: 2000-06-22
24 <160> NUMBER OF SEQ ID NOS: 28
26 <170> SOFTWARE: PatentIn version 3.0
28 <210> SEQ ID NO: 1
29 <211> LENGTH: 19
30 <212> TYPE: DNA
C--> 31 <213> ORGANISM: Artificial
33 <220> FEATURE:
34 <223> OTHER INFORMATION: sequence of a site in the PPAR response element
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37 tcaacacctta cccgtgtag                               19
40 <210> SEQ ID NO: 2
41 <211> LENGTH: 27
42 <212> TYPE: DNA
C--> 43 <213> ORGANISM: Artificial
45 <220> FEATURE:
46 <223> OTHER INFORMATION: primer
48 <400> SEQUENCE: 2
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52 <210> SEQ ID NO: 3
53 <211> LENGTH: 37
54 <212> TYPE: DNA
C--> 55 <213> ORGANISM: Artificial
57 <220> FEATURE:
58 <223> OTHER INFORMATION: primer
60 <400> SEQUENCE: 3
61 acgtgtcgac actagtggct agaggatctc taccagg           37
64 <210> SEQ ID NO: 4
65 <211> LENGTH: 48
66 <212> TYPE: DNA

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C--> 67 <213> ORGANISM: Artificial
      69 <220> FEATURE:
      70 <223> OTHER INFORMATION: primer
      72 <400> SEQUENCE: 4
      73 cgatgggtacc ctcgagcaat gtgctagcga gatccttcaa cctttacc      48
      76 <210> SEQ ID NO: 5
      77 <211> LENGTH: 13
      78 <212> TYPE: DNA
C--> 79 <213> ORGANISM: Artificial
      81 <220> FEATURE:
      82 <223> OTHER INFORMATION: sequence of site in PPAR response element
      84 <400> SEQUENCE: 5
      85 aggtcaaaagg tca      13
      88 <210> SEQ ID NO: 6
      89 <211> LENGTH: 69
      90 <212> TYPE: DNA
C--> 91 <213> ORGANISM: Artificial
      93 <220> FEATURE:
      94 <223> OTHER INFORMATION: primer
      96 <400> SEQUENCE: 6
      97 acgtgtcgac actaggtcaaa actaggtcaa aggtcacgga aaactaggtc aaaggtcacg      60
      99 gaaaactag      69
      102 <210> SEQ ID NO: 7
      103 <211> LENGTH: 64
      104 <212> TYPE: DNA
C--> 105 <213> ORGANISM: Artificial
      107 <220> FEATURE:
      108 <223> OTHER INFORMATION: primer
      110 <400> SEQUENCE: 7
      111 cgatgggtacc ctcgagcaat gtgctagcgg tgacctttga cctagtgtttc cgtgaccttt      60
      113 gacc      64
      116 <210> SEQ ID NO: 8
      117 <211> LENGTH: 32
      118 <212> TYPE: DNA
C--> 119 <213> ORGANISM: Artificial
      121 <220> FEATURE:
      122 <223> OTHER INFORMATION: primer
      124 <400> SEQUENCE: 8
      125 acgtagatct cgttagcggt gtacgggtggg ag      32
      128 <210> SEQ ID NO: 9
      129 <211> LENGTH: 29
      130 <212> TYPE: DNA
C--> 131 <213> ORGANISM: Artificial
      133 <220> FEATURE:
      134 <223> OTHER INFORMATION: primer
      136 <400> SEQUENCE: 9
      137 acgtaagctt ctatggaggt caaaacagc      29
      140 <210> SEQ ID NO: 10
      141 <211> LENGTH: 21

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142 <212> TYPE: DNA
C--> 143 <213> ORGANISM: Artificial
145 <220> FEATURE:
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148 <400> SEQUENCE: 10
149 ggtttgctga atgtgaagcc c                               21
152 <210> SEQ ID NO: 11
153 <211> LENGTH: 42
154 <212> TYPE: DNA
C--> 155 <213> ORGANISM: Artificial
157 <220> FEATURE:
158 <223> OTHER INFORMATION: primer
160 <400> SEQUENCE: 11
161 agtctctaga gctacgcgta caagtcttg tagatctcct gc       42
164 <210> SEQ ID NO: 12
165 <211> LENGTH: 32
166 <212> TYPE: DNA
C--> 167 <213> ORGANISM: Artificial
169 <220> FEATURE:
170 <223> OTHER INFORMATION: primer
172 <400> SEQUENCE: 12
173 agtcacgcgt gggcgatctt gacaggaaag ac                 32
176 <210> SEQ ID NO: 13
177 <211> LENGTH: 21
178 <212> TYPE: DNA
C--> 179 <213> ORGANISM: Artificial
181 <220> FEATURE:
182 <223> OTHER INFORMATION: primer
184 <400> SEQUENCE: 13
185 gcctttgagt gagctgatac c                               21
188 <210> SEQ ID NO: 14
189 <211> LENGTH: 35
190 <212> TYPE: DNA
C--> 191 <213> ORGANISM: Artificial
193 <220> FEATURE:
194 <223> OTHER INFORMATION: primer
196 <400> SEQUENCE: 14
197 agtcactagt aagctttttg ccgccagaac acagg              35
200 <210> SEQ ID NO: 15
201 <211> LENGTH: 36
202 <212> TYPE: DNA
C--> 203 <213> ORGANISM: Artificial
205 <220> FEATURE:
206 <223> OTHER INFORMATION: primer
208 <400> SEQUENCE: 15
209 agtcactagt ccatggctgc ccagtgcctc acgacc             36
212 <210> SEQ ID NO: 16
213 <211> LENGTH: 21
214 <212> TYPE: DNA

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C--> 215 <213> ORGANISM: Artificial
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      224 <210> SEQ ID NO: 17
      225 <211> LENGTH: 40
      226 <212> TYPE: DNA

C--> 227 <213> ORGANISM: Artificial
      229 <220> FEATURE:
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      232 <400> SEQUENCE: 17
      233 tgacgtgtcg acctagtaca agtccttgta gatctcctgc         40
      236 <210> SEQ ID NO: 18
      237 <211> LENGTH: 31
      238 <212> TYPE: DNA

C--> 239 <213> ORGANISM: Artificial
      241 <220> FEATURE:
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      244 <400> SEQUENCE: 18
      245 agtcgtcgac gcttcgagca gacatgataa g                 31
      248 <210> SEQ ID NO: 19
      249 <211> LENGTH: 35
      250 <212> TYPE: DNA

C--> 251 <213> ORGANISM: Artificial
      253 <220> FEATURE:
      254 <223> OTHER INFORMATION: primer
      256 <400> SEQUENCE: 19
      257 agtcgtcage gacggatcct tategatttt accac             35
      260 <210> SEQ ID NO: 20
      261 <211> LENGTH: 50
      262 <212> TYPE: DNA

C--> 263 <213> ORGANISM: Artificial
      265 <220> FEATURE:
      266 <223> OTHER INFORMATION: primer
      268 <400> SEQUENCE: 20
      269 gtcagctagc ctactcgagc caccatgggt gaaactctgg gagattctcc 50
      272 <210> SEQ ID NO: 21
      273 <211> LENGTH: 42
      274 <212> TYPE: DNA

C--> 275 <213> ORGANISM: Artificial
      277 <220> FEATURE:
      278 <223> OTHER INFORMATION: primer
      280 <400> SEQUENCE: 21
      281 tacggggtac ccagacatga taagatacat tgatgagttt gg       42
      284 <210> SEQ ID NO: 22
      285 <211> LENGTH: 33
      286 <212> TYPE: DNA

C--> 287 <213> ORGANISM: Artificial

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289 <220> FEATURE:
290 <223> OTHER INFORMATION: primer
292 <400> SEQUENCE: 22
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296 <210> SEQ ID NO: 23
297 <211> LENGTH: 33
298 <212> TYPE: DNA
C--> 299 <213> ORGANISM: Artificial
301 <220> FEATURE:
302 <223> OTHER INFORMATION: primer
304 <400> SEQUENCE: 23
305 tacgctcgag cttctatgga ggtcaaaaca gcg      33
308 <210> SEQ ID NO: 24
309 <211> LENGTH: 750
310 <212> TYPE: PRT
311 <213> ORGANISM: Homo sapiens
313 <400> SEQUENCE: 24
315 Met Gly Glu Thr Leu Gly Asp Ser Pro Ile Asp Pro Glu Ser Asp Ser
316 1 5 10 15
318 Phe Thr Asp Thr Leu Ser Ala Asn Ile Ser Gln Glu Met Thr Met Val
319 20 25 30
321 Asp Thr Glu Met Pro Phe Trp Pro Thr Asn Phe Gly Ile Ser Ser Val
322 35 40 45
324 Asp Leu Ser Val Met Glu Asp His Ser His Ser Phe Asp Ile Lys Pro
325 50 55 60
327 Phe Thr Thr Val Asp Phe Ser Ser Ile Ser Thr Pro His Tyr Glu Asp
328 65 70 75 80
330 Ile Pro Phe Thr Arg Thr Asp Pro Val Val Ala Asp Tyr Lys Tyr Asp
331 85 90 95
333 Leu Lys Leu Gln Glu Tyr Gln Ser Ala Ile Lys Val Glu Pro Ala Ser
334 100 105 110
336 Pro Pro Tyr Tyr Ser Glu Lys Thr Gln Leu Tyr Asn Lys Pro His Glu
337 115 120 125
339 Glu Pro Ser Asn Ser Leu Met Ala Ile Glu Cys Arg Val Cys Gly Asp
340 130 135 140
342 Lys Ala Ser Gly Phe His Tyr Gly Val His Ala Cys Glu Gly Cys Lys
343 145 150 155 160
345 Gly Phe Phe Arg Arg Thr Ile Arg Leu Lys Leu Ile Tyr Asp Arg Cys
346 165 170 175
348 Asp Leu Asn Cys Arg Ile His Lys Lys Ser Arg Asn Lys Cys Gln Tyr
349 180 185 190
351 Cys Arg Phe Gln Lys Cys Leu Ala Val Gly Met Ser His Asn Ala Ile
352 195 200 205
354 Arg Phe Gly Arg Met Pro Gln Ala Glu Lys Glu Lys Leu Leu Ala Glu
355 210 215 220
357 Ile Ser Ser Asp Ile Asp Gln Leu Asn Pro Glu Ser Ala Asp Leu Arg
358 225 230 235 240
360 Ala Leu Ala Lys His Leu Tyr Asp Ser Tyr Ile Lys Ser Phe Pro Leu
361 245 250 255

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RAW SEQUENCE LISTING ERROR SUMMARY      DATE: 05/21/2002  
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Input Set : A:\ST99021 Sequence.ST25.txt  
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Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,  
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23,26,27,28